



App No.:
Titl
Invemors:

09/693,205 Identification of ARSACS . . Thomas J. Hudson, *et al*.



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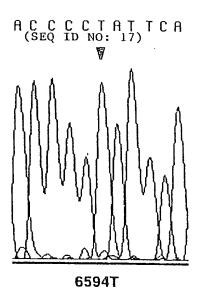
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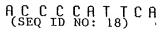
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APPROVED O.G. FIG.

BY CLASS SUBCLASS

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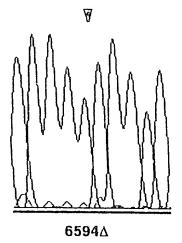


Figure 2A



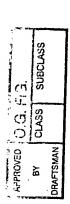
App No.: Title: Inventors: 09/693,205

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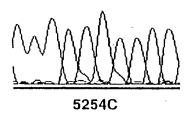
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A A A G C G A C A C (SEQ ID NO: 19)



A A A G M G A C A C (SEQ ID NO: 20)

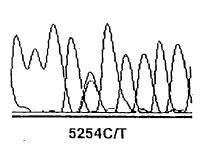


Figure 2B



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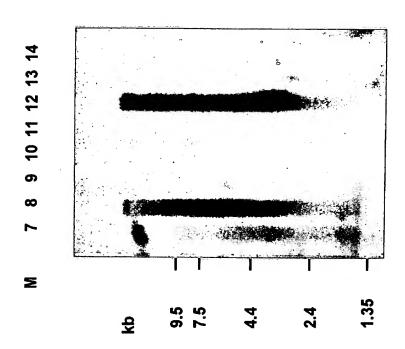
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APPROVED O.G. FIG.

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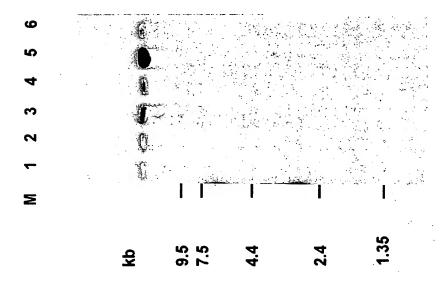


Figure 3

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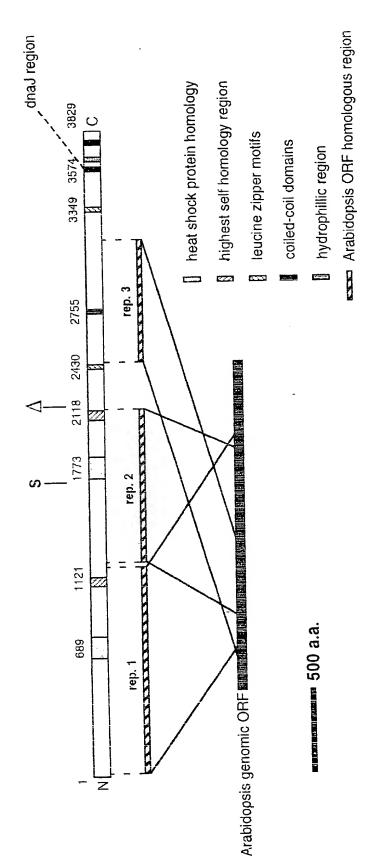


Figure 4A



SUBCLASS

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O.G. FIG. CLASS

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---EFMIMF<u>D</u>P

VYHITDIPIIMSR ... VYHITD.P. iS... $\frac{\text{VY}}{\text{If}}$.v.D.p..vs....

813 813 1879 1093

HSP90 (YEAST)

ARABIDOPSIS

1830 1044 78

HSP90 (YEAST)

HUMAN-REP 1 MOUSE-REP 1 MOUSE-REP 2

ARABIDOPSIS

764 764

HUMAN-REP 1 HUMAN-REP 1 HUMAN-REP 2

 $\dots E.ivMFDP$d.l.iFD.

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ATECSF <u>L</u> IDMRRNMDIRENLLDPG ATECSF <u>m</u> IDMRRNMDIRENLLDPG ATEF <u>v</u> .D.Rd.i.D ASE.tF <u>L</u> IDhd.LL.P.	ESLKRGE-VDKVGK <u>FGLGF</u> NS ESLKRGE-VDKVGK <u>FGLGF</u> NS K.GK.GK.G. <u>yGiGF</u> NS ga.KiGr <u>PGLG</u> FN. EalG.V.iG· <u>FGY</u> GF.S	Underline = conserved Bold = identical
GQREPLTVRIKNILEEYPSVSDIFKEILONADOAN-ATECSF <u>L</u> IDMRRNMDIRENLLDPG GQREPLTVRIKNILEEYPSVSDIFKEILONADOAN-ATECSF <u>M</u> IDMRRNMDIRENLLDPG GQKE.LT.RIK.IL. YPSem.KEILONADOAATEF <u>V</u> .D.Rd.i.D GQAE.LT.RIK.ILE.YIEI <u>V</u> ONAEOAASE.tF <u>L</u> IDhd.LL.P.	MAACHGPALWSF <u>N</u> NSQ-F <u>SDSD</u> FVNITRLGESLKRGE-VDKVGK <u>FGLGF</u> NS MAACHGPALWSF <u>N</u> NSG-F <u>SDSD</u> FINITRLGESLKRGE-VDKVGK <u>FGLGF</u> NS .AGPALY <u>N</u> NF <u>L</u> E. <u>D</u> I.LGK.GK.GK.G. MAGPALY.F <u>N</u> NSF <u>L</u> DISRİGqa.KİG <u>F<u>P</u>GLGFN.</u>	<u>VY</u> HITDIPIIMSREFMIMF <u>D</u> P VYHITDIPIIMSREFMIMF <u>D</u> P

705 705 1773 987 18

HSP90 (YEAST)

ARABIDOPSIS

HUMAN-REP 2

MOUSE-REP 1

Figure 4B

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APPROVED O.G. FIG.
BY CLASS SU

MNTFWPGRELIVQWYPFDENRNHPSVSWLKMVWKNLYIHFSEDLTLFDEMPLIPRTILEEVS.DKRLL.N.	60
GQTCVELIRLRIPSLVILDDESEAQLPEFLADIVQKLGGFVLKKLDASIQHPLIKKYIHS DVTIRTVV	120
PLPSAVLQIMEKMPLQKLCNQITSLLPTHKDALRKFLASLTDSSEKEKRIIQELAIFKRI	180
NHSSDQGISSYTKLKGCKVLHHTAKLPADLRLSISVIDSSDEATIRLANMLKIEQLKTTS	240
CLKLVLKDIENAFYSHEEVTQLMLWVLENLSSLKNENPNVLEWLTPLKFIQISQEQMVSAFGTQISDMHMGHV.A.	300
GELFDPDIEVLKDLFCNEEGTYFPPSVFTSPDILHSLRQIGLKNEASLKEKDVVQVAKKI.DRYEACTISR	360
EALQVGACPDQDVLLKKAKTLLLVLNKNHTLLQSSEGKMTLKKIKWVPACKERPPNYPGSSS.QNMQAA	420
LVWKGDLCNLCAPPDMCDVGHAILIGSSLPLVESIHVNLEKALGIFTKPSLSAVLKHFKIAAV.VVQSTINT	480
VVDWYSSKTFSDEDYYQFQHILLEIYGFMHDHLNEGKDSFRALKFPWVWTGKKFCPLAQATNN	540
VIKPIHDLDLQPYLHNVPKTMAKFHQLFKVCGSIEELTSDHISMVIQKIYLKSDQDLSEQTYAVEE	600
ESKQNLHLMLNIIRWLYSNQIPASPNTPVPIHHSKNPSKLIMKPIHECCYCDIKVDDLND	660
LLEDSVEPIILVHEDIPMKTAEWLKVPC LSTRLINPENMGFEQSGQREPLTVRIKNILEE	720
YPSVSDIFKELLONADDANATECSFLIDMRRNMDIRENLLDPGMAACHGPALWSFNNSQF	780
YPSVSDIFKELLQNADDANATECSFLIDMRRNMDIRENLLDPGMAACHGPALWSFNNSQFE.	780
	780 840
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISK	
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISK LL HIKDKSNPGIKINWSKQQKRLRKFPNQFKPFIDVFGCQLPLTVEAPYSYNGTLFRLSFRT	840
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISK LL HIKDKSNPGIKINWSKQQKRLRKFPNQFKPFIDVFGCQLPLTVEAPYSYNGTLFRLSFRTRAA	840 900
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISK L HIKDKSNPGIKINWSKQQKRLRKFPNQFKPFIDVFGCQLPLTVEAPYSYNGTLFRLSFRT R. A	840 900 960
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISK L HIKDKSNPGIKINWSKQQKRLRKFPNQFKPFIDVFGCQLPLTVEAPYSYNGTLFRLSFRT R. A QQEAKVSEVSSTCYNTADIYSLVDEFSLCGHRLIIFTQSVKSMYLKYLKIEETNPSLAQD	840 900 960 1020
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISKL HIKDKSNPGIKINWSKQQKRLRKFPNQFKPFIDVFGCQLPLTVEAPYSYNGTLFRLSFRTR. QQEAKVSEVSSTCYNTADIYSLVDEFSLCGHRLIIFTQSVKSMYLKYLKIEETNPSLAQDN TVIIKKKSCSSKALNTPVLSVLKEAAKLMKTCSSSNKKLPSDEPKSSCILQITVEEFHHV .IV.PA FRRIADLQSPLFRGPDDDPAALFEMAKSGQSKKPSDELSQKTVECTTWLLCTCMDTGEALT. KFSLSESGRRLGLVPCGAVGVQLSEIQDQKWTVKPHIGEVFCYLPLRIKTGLPVHINGCF	840 900 960 1020 1080
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISK L HIKDKSNPGIKINWSKQQKRLRKFPNQFKPFIDVFGCQLPLTVEAPYSYNGTLFRLSFRT R. A	840 900 960 1020 1080
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISKL	840 900 960 1020 1080 1140
SDSDFVNITRLGESLKRGEVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISK LL HIKDKSNPGIKINWSKQQKRLRKFPNQFKPFIDVFGCQLPLTVEAPYSYNGTLFRLSFRTRA QQEAKVSEVSSTCYNTADIYSLVDEFSLCGHRLIIFTQSVKSMYLKYLKIEETNPSLAQDN TVIIKKKSCSSKALNTPVLSVLKEAAKLMKTCSSSNKKLPSDEPKSSCILQITVEEFHHV .IV.PAT.V FRRIADLQSPLFRGPDDDPAALFEMAKSGQSKKPSDELSQKTVECTTWLLCTCMDTGEALTPDI KFSLSESGRRLGLVPCGAVGVQLSEIQDQKWTVKPHIGEVFCYLPLRIKTGLPVHINGCFNL.H.T.EI AVTSNRKEIWKTDTKGRWNTTFMRHVIVKAYLQVLSVLRDLATSGELMDYTYYAVWPDPDAIGT LVHDDFSVICQGFYEDIAHGKGKELTKVFSDGSTWVSMKNVRFLDDSILKRRDVGSAAFKKRMQ.K	840 900 960 1020 1080 1140 1200 1260

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PPROVED	0.G. FIG.	-IG.
ВУ	CLASS	SUBCLASS
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LIDEKLKIRDPRAKDFAAKYQTIRFLPFLTKPAGFSLDWKGNSFKPETMFAATDLYTAEH	1500
QDIVCLLQPILNENSHSFRGCGSVSLAVKEFLGLLKKPTVDLVINQLKEVAKSVDDGITLQQ	1560
YQENITNACYKYLHEALMQNEITKMSIIDKLKPFSFILVENAYVDSEKVSFHLNFEAAPYVLMA.ATECVE	1620
LYQLPNKYKNNFRELFETVGVRQSCTVEDFALVLESIDQERGTKQITEENFQLCRRIISEKK	1680
GIWSLIREKKQEFCEKNYGKILLPDTNLMLLPAKSLCYNDCPWIKVKDTTVKYCHADIPRRSS	1740
EVAVKLGAVPKRHKALERYASNVCF TTLGTEFGQKEKLTSRIKSILNAYPSEKEMLKELL I	1800
QNADDAKATEICFVFDPRQHPVDRIFDDKWAPLQGPALCVYNNQPFTEDDVRGIQNLGKG	1860
TKEGNPYKTGQYGIGFNSVYHITDCPSFISGNDILCIFDPHARYAPGATSISPGRMFRDLGHVVV	1920
DADFRTQFSDVLDLYLGTHFKLDNCTMFRFPLRNAEMAKVSEISSVPASDRMVQNLLDKL	1980
RSDGAELLMFLNHMEKISICEIDKSTGALNVLYSVKGKITDGDRLKRKQFHASVIDSVTKAGAG	2040
KRQLKDIPVQQITYTMDTEDSEGNLTTWLICNRSGFSSMEKVSKSVISAHKNQDITLFPR	2100
GGVAACITHNYKKPHRAFCFLPLSLETGLPFHVNGHFALDSARRNLWRDDNGVGVRSDWN	2160
NSLMTALIAPAYVELLIQLKKRYFPGSDPTLSVLQNTPIHVVKDTLKKFLSFFPVNRLDL	2220
QPDLYCLVKALYNCIHEDMKRLLPVVRAPNIDGSDLHSAVIITWINMSTSNKTRPFFDNL	2280
LQDELQHLKNADYNITTRKTVAENVYRLKHLLLEIGFNLVYNCDETANLYHCLIDADIPV	2340
SYVTPADIRSFLMTFSSPDTNCHIGKLPCRLQQTNLKLFHSLKLLVDYCFKDAEENEIEVVS.F	2400
EGLPLLITLDSVLQTFDAKRPKFLTTYHELIPSRKDLFMNTLYLKYSNILLNCKVAKVFD	2460
ISSFADLLSSVLPREYKTKSCTKWKDNFASESWLKNAWHFISESVSVKEDQEETKPTFDITDPAV	2520
VVDTLKDWALLPGTKFTVSANQLVVPEGDVLLPLSLMHIAVFPNAQSDKVFHALMKAGCI I.ITSI	2580
QLALNKICSKDSAFVPLLSCHTANIESPTSILKALHYMVQTSTFRAEKLVENDFEALLMY	2640
FNCNLNHLMSQDDIKILKSLPCYKSISGRYVSIGKFGTCYVLTKSIPSAEVEKWTQSSSS	2700
AFLEEKIHLKELYEVIGCVPVDDLEVYLKHLLPKIENLSYDAKLEHLIYLKNRL <u>SSAEEL</u> V	2760
SEIKEQLFEKLESLLIIHDANSRLKQAKHFYDRTVRVFEVMLPEKLFIPNDFFKKLEQLI	2820
KPKNHVTFMTSWVEFLRNIGLKYILSQQQLLQFAKEISVRANTENWSKETLQNTVDILLH QAA ASS	2880

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O.G. FIG.	CLASS SUBCLASS	
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HIFQERMDLLSGNFLKELSLIPFLCPERAPAEFIRFHPQYQEVNGTLPLIKFNGAQVNPK	2940
FKQCDVLQLLWTSCPILPEKATPLSIKEQEGSDLGPQEQLEQVLNMLNVNLDPPLDKVINAA	3000
NCRNICNITTLDEEMVKTRAKVLRSIYEFLSAEKREFRFQLRGVAFVMVEDGWKLLKPEE	3060
VVINLEYESDFKPYLYKLPLELGTFHQLFKHLGTEDIISTKQYVEVLSRIFKNSEGKQLDASS	3120
PNEMRTVKRVVSGLFRSLQNDSVKVRSDLENVRDLALYLPSQDGRLVKSSILVFDDAPHY	3180
KSRIQGNIGVQMLVDLSQCYLGKDHGFHTKLIMLFPQKLRPRLLSSILEEQLDEETPKVC	3240
QFGALCSLQGRLQLLLSSEQFITGLIRIMKHENDNAFLANEEKAIRLCKALREGLKVSCF	3300
EKLQTTLRVKGFNPIPHSRSETFAFLKRFGNAVILLYIQHSDSKDINFLLALAMTLKSAT	3360
DNLISDTSYLIAMLGCNDIYRIGEKLDSLGVKYDSSEPSKLELPMPGTPIPAEIHYTLLM	3420
DPMNVFYPGEYVGYLVDAEGGDIYGSYQPTYTYAIIVQEVEREDADNSSFLGKIYQIDIG	3480
YSEYKIVSSLDLYKFSRPEESSQSRDSAPSTPTSPTEFLTPGLRSIPPLFSGRESHKT-S	3540
SKHQSPKKLKVNSLPEILKEVTSVVEQAWKL <i>PESERKKIIRRLYLKWHPDKNPFNHDIAN</i> T.H.RA	3600
<u>EVFKHLONEINRLEKOA</u> FLDQNADRASRRTFSTSASRFQSDKYSFQRFYT <u>SWNQEATSHK</u>	3660
SERQQQNKEKCPPSAGQTYSQRFFVPPTFKSVGNPVEARRWLRQARANFSAARNDLHKNA	3720
NEWVCFKCYLSTKLALIAADYAVRGKSDKDVKP <u>TALAQKIEEYSQQLEGLTNDVHTLEAY</u>	3780
GVDSLKTRYPDLLPFPQIPNDRFTSEVAMRVMECTACIIIKLENFMQQKV	3830



SUBCLASS

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APPROVED O.G. FIG.

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Table 1 ESTs identified by sample-sequencing of the ARSACS critical interval

BAC clone	GenBank #	UniGene	Identity*	Tissue Source
235_1,_20	AA987300	IIs. 129092	221/230	neuroendocrine lung carcinoids
235_1,_20	AA476635*		249/296	total fetus
235_1,_20	AI351876*		272/335	melanocyte, fetal heart, pregnant uterus (pool)
235_1,_20	W25994	IIs. 163732	447/464	retina
235_1,_20	AI377467	IIs. 163732	257/263	total fetus
235_1,_20	AA601007		488/491	schwannoma tumor
235_1,_20	AA324964		214/234	cerebellum
235_1,_20, 206_1,_I	AA897178 ^b		238/238	fetal lung, testis, and B-cell
235_1,_20, 206_1,_1	$RI7106^{b}$	IIs. 188560	747/784	brain, adipose tissue
235_I,_20, 206_I,_I	AB018273 ^b	IIs. 159492	4318/4318	multiple tissue types including brain, CNS, and whole embryo

Number of homologous nucleotides between the BAC subclone sequence and the GenBank sequence.

ж. ф. *.

AA897178, RI7106, and AB018273 are all contained within *spastin*. AA176335 and AI351876 were homologous to non-overlapping portions of the same M13 subclone sequence.

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Table 2 Primers for PCR amplification of the human spastin UKF

Product size (bp)

Reverse primer

Forward primer

Primer set

	APPROVED O.G. FIG.	O.G. F	<u>1</u> 6.
	β	CLASS	CLASS SUBCLASS
	DRAFTSMAN		
Poblo 2 Drimore for DCD amplification of the human enactin ODE	nactive C	NDE	

603	581	592	602	611	624	631	620	604	651	625	979	624	648	609	631	609	645	618	612	647	654	638
(SEQ ID NO: 22)	(SEQ ID NO: 24)	(SEQ ID NO: 26)	(SEQ ID NO: 28)	(SEQ ID NO: 30)	(SEQ ID NO: 32)	(SEQ ID NO: 34)	(SEQ ID NO: 36)	(SEQ ID NO: 38)	(SEQ ID NO: 40)	(SEQ ID NO: 42)	(SEQ ID NO: 44)	(SEQ ID NO: 46)	(SEQ ID NO: 48)	(SEQ ID NO: 50)	(SEQ ID NO: 52)	(SEQ ID NO: 54)	(SEQ ID NO: 56)	(SEQ ID NO: 58)	(SEQ ID NO: 60)	(SEQ ID NO: 62)	(SEQ ID NO: 64)	(SEQ ID NO: 66)
CAAGAACTTCCTCAGGGCATC	GGTGGGAAATAGGTTCCTTC	GCACTAAGGCTAGGTTTTGTGAAG	CGTGAATTGGCTTCATGATAA	GATGGGAATGTCAGTGATATGG	CTTTGGTTCATCACTGGGAAG	CAGGTCCCGTAAGACACTCAG	CGAAGAACTCCCGAGAACTCA	GCAAACATGGTTTCAGGCTTA	ATTATTCGTCGCCAAAGCTGA	ACACAAAGTGCTGGCCCTTGC	ATACAGCACATTTAGAGCTCCAGT	GCAATTCAACATATGCAGGAG	TGATATCAGCAGGGTCACAT	GCCATGCATTCTTAAGCCAAG	AGCGCCACTGATGGATTTAT	TTCCACCCAGGATGTCATAAA	ATCAAGAGGAGGATCCAGGTT	TAAAGCGCAAGGTCTCGTACA	TCTGCTGTGGGGAATAGGATT	TGCTTTGAGAGCTTTCCTCAG	GTAAGTCTGTCCGGCTGAAGG	TTCGTGCTACACACATTCAAGA
CCTTCCAGTACTGTGTTATTTGTGAG	GATGCATCTATACAACATCCGCT	AAAAATGAGAATCCAAATGTGCT	GCTCCTCACTTCCTCTTGTTG	AGCAATCAGATTCCAGCAAGC	GGGAGAAGTTGACAAAGTTGGA	TCCAAAGCATTGAACACCCT	CAATGGGTGCTTTGCTGTTAC	GCTGCCTAAACAGATACTAC	CAAACAATCCGCTTCCTTCCAT	TTCCGCGAACTTTTTGAAACC	GATGCAAAGGCGACAGAAATC	GCATCAGACAGAATGGTCCAG	GTGAATGGCCACTTTGCACT	ACCACACGCAAAACAGTAGCA	TGACATTTCCAGCTTTGCTGA	AAATGATTTTGAGGCACTTTTG	ACAGTAGACTAAAGCAAGCAAAGC	CATCCTGCCCTATTCTTCCAG	TGAGGCAAACAATTAGATCC	GCAAAGCCCTAAGAGAAGGATT	TGAAAGAGAAGATGCTGACAATTC	CATCCCGATTTCAGTCAGACA
(SEQ ID NO: 21)	(SEQ ID NO: 23)	(SEQ ID NO: 25)	(SEQ ID NO: 27)	(SEQ ID NO: 29)	(SEQ ID NO: 31)	(SEQ ID NO: 33)	(SEQ ID NO: 35)	(SEQ ID NO: 37)	(SEQ ID NO: 39)	11 (SEQ ID NO: 41)	(SEQ ID NO: 43)	(SEQ ID NO: 45)	(SEQ ID NO: 47)	(SEQ ID NO: 49)	16 (SEQ ID NO: 51)	(SEQ ID NO: 53)	(SEQ ID NO: 55)	19 (SEQ ID NO: 57)	(SEQ ID NO: 59)	21 (SEQ ID NO: 61)	(SEQ ID NO: 63)	(SEQ ID NO: 66)
	2	3	4	5	9	7	8	6	10	11	12	13	14	15	16	17	18	19	20	21	22	23



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SUBCLASS E G 0.0 CLASS **APPROVED** ₽

```
ROD
                         11493 bp
                                      DNA
LOCUS
            AF193557
DEFINITION
            Mus musculus sacsin gene, complete cds.
ACCESSION
            AF193557
            AF193557.1
                       GI:6907043
VERSION
KEYWORDS
SOURCE
            house mouse.
            Mus musculus
  ORGANISM
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
REFERENCE
               (bases 1 to 11493)
            Engert, J.C., Berube, P., Mercier, J., Dore, C., Lepage, P., Ge, B.,
  AUTHORS
            Bouchard, J.P., Mathieu, J., Melancon, S.B., Schalling, M.,
            Lander, E.S., Morgan, K., Hudson, T.J. and Richter, A.
            ARSACS, a spastic ataxia common in northeastern Quebec, is caused
  TITLE
            by mutations in a new gene encoding an 11.5-kb ORF
            Nat. Genet. 24 (2), 120-125 (2000)
  JOURNAL
  MEDLINE
            20120709
REFERENCE
                (bases 1 to 11493)
            Engert, J.C., Berube, P., Dore, C., Lepage, P., Ge, B., Hudson, T.J. and
  AUTHORS
            Richter, A.
            Direct Submission
  TITLE
            Submitted (08-OCT-1999) Genome Centre, Montreal General Hospital,
  JOURNAL
            1650 Cedar Ave., Montreal, QC H3G 1A4, Canada
FEATURES
                     Location/Qualifiers
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                      /db_xref="taxon:10090"
                      <1..>11493
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                      /product="sacsin"
     CDS
                      1..11493
                      /note="molecular chaperone"
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                      /product="sacsin"
                      /protein_id="AAF31263.1"
                      /db xref="GI:6907044"
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SUBCLASS O.G. FIG. CLASS

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THE WATER NTPVPIYHSRNPSKLVMKPIHECCYCDIKVDDLNDLLEDSVEPIILVHEDIPMKTAE WLKVPCLSTRLINPENMGFEQSGQREPLTVRIKNILEEYPSVSDIFKELLQNADDANA TECSFMIDMRRNMDIRENLLDPGMAACHGPALWSFNNSEFSDSDFLNITRLGESLKRG EVDKVGKFGLGFNSVYHITDIPIIMSREFMIMFDPNINHISKHIKDRSNPGIKINWSK OOKRLRKFPNOFKPFIDVFGCQLPLAVEAPYSYNGTLFRLSFRTQQEAKVSEVSSTCY NTADIYSLVDEFSLCGHRLIIFTQSVNSMYLKYLKIEETNPSLAQDTIIIKKKVCPSK ALNAPVLSVLKEAAKLMKTCSSSNKKLPTDVPKSSCILQITVEEFHHVFRRIADLQSP LFRGPDDDPATLFEMAKSGQSKKPSDELPQKTVDCTTWLICTCMDTGEALKFSLNESG RRLGLVPCGAVGVLLHETQEQKWTVKPHIGEVFCYLPLRIKTGLPIHINGCFAVTSNR KEIWKTDTKGRWNTTFMRHVIVKAYLQALSVLRDLAIGGELTDYTYYAVWPDPDLVHD DFSVICKGFYEDIAHGKGKELTRVFSDGSMWVSMKNVRFLDDSILQRKDVGSAAFKIF LKYLKKTGSKNLCAVELPSSVKAGFEEAGCKQILLENTFSEKQFFSEVFFPNIQEIEA ELRDPLMNFVLNEKLDEFSGILRVTPCVPCSLEGHPLVLPSRLIHPEGRVAKLFDTKD GRFPYGSTODYLNPIILIKLVQLGMAKDDILWDDMLERAESVAEINKSDHAAACLRSS ILLSLIDEKLKIKDPRAKDFAAKYQTIPFLPFLTKPAGFSLEWKGNSFKPETMFAATD $\verb|IYTAEYQDIVCLLQPILNENSHSFRGCGSVSLAVKEFLGLLKKPTVDLVINQLKQVAK|$ SVDDGITLYQENITNACYKYLHEAVLQNEMAKATIIEKLKPFCFILVENVYVESEKVS FHLNFEAAPYLYQLPNKYKNNFRELFESVGVRQSFTVEDFALVLESIDQERGKKQITE ENFQLCRRIISEGIWSLIREKRQEFCEKNYGKILLPDTNLLLLPAKSLCYNDCPWIKV KDSTVKYCHADIPREVAVKLGAIPKRHKALERYASNICFTALGTEFGQKEKLTSRIKS ILNAYPSEKEMLKELLQNADDAKATEICFVFDPRQHPVDRIFDDKWAPLQGPALCVYN NOPFTEDDVRGIQNLGKGTKEGNPCKTGHYGIGFNSVYHITDCPSFISGNDILGIFDP HARYAPGATSVSPGRMFRDLDADFRTQFSDVLDLYLGNHFKLDNCTMFRFPLRNAEMA QVSEISSVPSSDRMVQNLLDKLRSDGAELLMFLNHMEKISICEIDKATGGLNVLYSVK GKITDGDRLKRKQFHASVIDSVTKKRQLKDIPVQQITYTMDTEDSEGNLTTWLICNRS GFSSMEKVSKSVISAHKNQDITLFPRGGVAACITHNYKKPHRAFCFLPLSLETGLPFH VNGHFALDSARRNLWRDDNGVGVRSDWNNSLMTALIAPAYVELLIQLKKRYFPGSDPT LSVLQNTPIHVVKDTLKKFLSFFPVNRLDLQPDLYCLVKALYSCIHEDMKRLLPVVRA

Figure 8B



SUBCLASS

O.G. FIG.

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PNIDGSDLHSAVIITWINMSTSNKTRPFFDNLLQDELQHLKNADYNITTRKTVAENVY RLKHLLLEIGFNLVYNCDETANLYHCLVDADIPVSYVTPADVRSFLMTFSSPDTNCHI GKLPCRLQQTNLKLFHSLKLLVDYCFKDAEESEFEVEGLPLLITLDSVLQIFDGKRPK FLTTYHELIPSRKDLFMNTLYLKYSSVLLNCKVAKVFDISSFADLLSSVLPREYKTKN CAKWKDNFASESWLKNAWHFISESVSVTDDQEEPKPAFDVIVDILKDWALLPGTKFTV STSOLVVPEGDVLIPLSLMHIAVFPNAQSDKVFHALMKAGCIQLALNKICSKDSALVP LLSCHTANIDSPASILKAVHYMVQTSTFRTEKLMENDFEALLMYFNCNLSHLMSQDDI KILKSLPCYKSISGRYMSIAKFGTCYVLTKSIPSAEVEKWTQSSSSAFLEEKVHLKEL YEVLGCVPVDDLEVYLKHLLPKIENLSYDAKLEHLIYLKNRLASIEEPSEIKEQLFEK LESLLIIHDANNRLKQAKHFYDRTVRVFEVMLPEKLFIPKEFFKKLEQVIKPKNQAAF MTSWVEFLRNIGLKYALSQQQLLQFAKEISVRANTENWSKETLQSTVDILLHHIFQER MDLLSGNFLKELSLIPFLCPERAPAEYIRFHPQYQEVNGTLPLIKFNGAQVNPKFKQC DVLQLLWTSCPILPEKATPLSIKEQEGSDLAPQEQLEQVLNMLNVNLDPPLDKVINNC RNICNITTLDEEMVKTRAKVLRSIYEFLSAEKREFRFQLRGVAFVMVEDGWKLLKPEE VVINLEYEADFKPYLYKLPLELGTFHQLFKHLGTEDIISTKQYVEVLSRIFKSSEGKQ $\verb|LDPNEMRTVKRVVSGLFKSLQNDSVKVRSDLENARDLALYLPSQDGKLVKSSILVFDD|$ APHYKSRIQGNIGVQMLVDLSQCYLGKDHGFHTKLIMLFPQKLRPRLLSSILEEQLDE ETPKVCQFGALCSLQGRLQLLLSSEQFITGLIRIMKHENDNAFLANEEKAIRLCKALR EGLKVSCFEKLQTTLRVKGFNPIPHSRSETFAFLKRFGNAVILLYIQHSDSKDINFLL ALAMTLKSATDNLISDTSYLIAMLGCNDIYRISEKLDSLGVKYDSSEPSKLELPMPGT PIPAEIHYTLLMDPMNVFYPGEYVGYLVDAEGGDIYGSYQPTYTYAIIVQEVEREDAD NTSFLGKIYQIDIGYSEYKIVSSLDLYKFSRPDESSQNRDSAPTTPTSPTEFLTPGLR SIPPLFSGKESHKSPSTKHHSPRKLKVNALPEILKEVTSVVEQAWKLPESERKKIIRR LYLKWHPDKNPENHDIANEVFKHLQNEINRLEKQAFLDQNADRASRRTFSTSASRFQS DKYSFQRFYTSWNQEATSHKSERQQQSKEKCPPSAGQTYSQRFFVPPTFKSVGNPVEA RRWLRQARANFSAARNDLHKNANEWVCFKCYLSTKLALIAADYAVRGKSDKDVKPTAL AQKIEEYSQQLEGLTNDVHTLEAYGVDSLKTRYPDLLPFPQIPNDRFTSEVAMRVMEC TACIIIKLENFIQQKV"

Figure 8C

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BASE COUNT 3599 a 2281 c 2387 g 3226 t ORIGIN

						
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					agaatctcta	
					ctagaactct	
181	gaccagacgt	gtgtggaact	catcagactc	aggatcccat	cagtagtcat	tttagatgat
					tacaaaaact	
					ttaaaaaata	
361	ccactcccga	gtgctatttt	gcagataatg	gagaagatac	ctctacagaa	gttgtgtaat
					ggaagttttt	
481	actgatacca	gtgaaaaaga	gaaaagaata	attcaagaat	tgacaatatt	caaaagaatt
541	aatcactcat	cagatcaagg	gatttcctct	tacacaaaat	taaaaggatg	taaagttttg
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841	cttgattggt	taatgccact	aaaattcatt	catatgtccc	agggacatgt	ggtagcagct
901	ggtgatctct	ttgatcctga	tatagaagta	ctaagggatc	tcttttataa	tgaagaagaa
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1021	ggcttaaaaa	atgaatccag	tctaaaagaa	aaagatgttg	tacaagtggc	aagaaaaatt
1081	gaagctttac	aggtcagttc	ctgtcagaat	caggatgttc	tcatgaagaa	agccaaaaca
1141	ctcttactgg	tcttgaataa	aaaccagaca	ctcttgcagt	cttctgaagg	gaagatggca
					ctccaaatta	
1261	ttagtctgga	aaggggatct	ctgtaatctt	tgtgcacctc	cagatatgtg	tgatgcggca
1321	catgcagttc	tagtaggctc	ctcacttcct	cttgttgaaa	gtgtccatgt	gaacctggag
1381	caggcgctca	gcatcttcac	aaagcctact	atcaatgctg	tcttaaaaca	ctttaaaact
1441	gttgttgact	ggtatacttc	aaaaaccttt	agtgatgaag	attactatca	gttccaacat
1501	attttgcttg	aaatttatgg	gttcatgcat	gatcatctga	gtgaagggaa	ggattctttt
1561	aaagccttga	agtttccatg	ggtttggact	ggcaaaaact	tttgtcctct	tgcccaggct
					tatataatgt	
1681	atggcaaaat	tccaccagct	gttcaaggct	tgtggctcaa	tagaagagtt	gacatcagat
1741	catatttcca	tggtcattca	gaaagtttat	ctcaaaagtg	accaggagtt	gagtgaagaa
1801	gaaagtaaac	aaaatcttca	tctcatgttg	aatattatga	gatggctcta	tagcaatcag
					gcagaaatcc	
					aagttgatga	
1981	ttgcttgaag	attcagtgga	accaattatc	ttggtacatg	aagatatacc	catgaaaact
2041	gcagaatggc	taaaagttcc	gtgccttagt	acaagactga	tcaatcctga	aaacatgggg
					ttaaaaatat	
					atgctgatga	
					acatacggga	
					cattcaacaa	
					ctttaaaaag	
					accacatcac	
					acataaatca	
					ggagtaagca	
2561	ttaaggaagt	nagataatta	guttaaacta	gazatatt	tatttggctg	ctttagecacce
					tccgactgtc	
					acaatactgc tcatttttac	
2/01	anatagatat	atttassets	attananatt	cacagaccca	atectacett	accacaacat
2021	accepacyc	ttaagaaaaa	agtttgggg	tccaaaccat	atcctagctt tgaatgcacc	agttttaagt
					gcagcaacaa	
					tcgaagaatt	
					gtccagatga	
					catcagatga	
					tggatacagg	
					ttccttgtgg	
2241	~~g	-juu-guudg				JJJJ-5



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Inventors:

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APPROVED O.G. FIG.

BY CLASS SUBCLASS

FRAFTSMAN

3301	gttctcttgc	atgaaaccca	ggaacagaag	tggaccgtga	aaccacacat	aggagaagtg
3361	ttttgctatt	tacctctacg	aatcaaaaca	gggttgccaa	ttcacatcaa	tgggtgcttt
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3481	acattcatga	ggcatgtcat	tgtgaaagct	tacttacaag	ccctcagtgt	cttacgggac
3541	ctagccattg	gtggtgagct	gactgattat	acttactatg	cagtgtggcc	tgatcctgat
3601	ctagttcatg	atgacttctc	tgtgatctgt	aaaggatttt	atgaagacat	tgctcatggg
3661	aaggggaagg	agttgaccag	agtcttctct	gatgggtcta	tgtgggtttc	catgaagaat
3721	gtgaggtttc	tggatgactc	tatacttcaa	aggaaagatg	ttggttcagc	agccttcaag
					tctgtgctgt	
					agatactgct	
3901	ttttcagaga	aacagttctt	ttcagaagtc	ttctttccta	atatccagga	aattgaagca
3961	gaacttagag	atcctctgat	gaattttgtc	ctaaatgaaa	aacttgatga	gttctcagga
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					ggagtagtat	
					aggattttgc	
					gtttttcttt	
					acatttacac	
					attcccattc	
4561	tgtggttcag	tgtctttggc	tgttaaggag	tttttgggtt	tactaaagaa	gccaacagtt
4621	qatctqqtaa	taaaccaqtt	qaagcaagtt	gcaaaatcag	ttgatgatgg	cattacattg
					atgaagcagt	
					tttgtttcat	
					actttgaagc	
					agctttttga	
4921	gtgcgacagt	catttactqt	tgaagacttt	gccctagttt	tggagtctat	tgatcaagag
4981	agaggaaaaa	aacaaataac	agaagagaat	tttcagcttt	gccgacgaat	aatcagtgaa
5041	ggcatctgga	gtctcattag	agaaaagaga	caaqaatttt	gtgagaaaaa	ttatggcaaa
5101	atattactqc	cagacactaa	cctqctqctq	ctccctgcta	agtcattatg	ctacaatgac
5161	tgtccctgga	taaaagtaaa	ggactccact	gtcaagtatt	gccatgccga	cataccccgg
5221	gaagtagctg	taaaacttgg	tgcaatacca	aagagacata	aagcattaga	aagatatgca
5281	tccaacatct	gtttcacagc	tctaggtaca	gaatttgggc	agaaagaaaa	actgaccagc
					aaatgctgaa	
					tgtttgatcc	
					aagggccagc	
					ttcagaatct	
					tcggattcaa	
5641	catattacag	actgcccttc	ttttatttct	ggcaatgaca	tcctgggtat	ttttgatccc
					gacgcatgtt	
5761	gatgcagact	ttagaaccca	gttctcagat	gttctagatc	tgtacttggg	aaaccacttt
5821	aaactggaca	attgtacaat	gtttagattt	cctctgcgta	atgcagagat	ggcacaagtt
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APPROVED O.G. FIG.

BY CLASS SUBCLASS

DRAFTSMAN

6721	cgtcttttgc	ctgttgttcg	ggctccaaat	attgatggct	cagatttgca	ctctgcagtc
6781	ataattactt	ggatcaatat	gtctacttca	aataaaacta	gaccattttt	tgataactta
6841	ctacaggatg	aattacagca	ccttaaaaat	gcagattata	acatcacaac	tcgaaaaaca
6901	gtcgcagaga	atgtctacag	actgaagcac	ctgctcttag	aaattggttt	caacttggtt
6961	tataactgtg	atgaaactgc	taacctttac	cattgccttg	tagatgcaga	tatccctgtc
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7081	aattgccata	ttgggaagct	gccttgtcgt	cttcagcaga	ctaacctaaa	actttttcac
7141	agtttaaaac	ttttagttga	ttactgtttt	aaagatgctg	aagaaagtga	gtttgaagtt
7201	gagggactgc	ccctactcat	tacactggac	agtgtcttgc	agatttttga	tggtaaacga
7261	cccaagtttc	taacaacata	ccatgaatta	attccatcgc	gtaaagactt	gtttatgaac
7321	accttatact	tgaaatacag	tagtgttttg	ttgaactgca	aagttgcaaa	agtgtttgac
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	atcagtgaat					
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	agtcagcttg					
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	gatgcaaagt					
	tcagagatta					
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	aaggtcctaa					
	cttcggggtg					
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	aaaagtagga					
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	cctcgtctgc					
	cagtttggcg					
	ttcatcacag					
	gaagaaaaag					
	gagaagcttc					
	gaaactttcg					
	tcagacagca					
	gacaatttga					
	Jacancesgu		Joannacoca		- 3 3 3 - 0 3 0 4 4	-3



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10141 aggatcagtg agaagcttga cagtttaggg gtgaaatacg actcctctga gccatcaaaa 10201 ctggaactcc ccatgcctgg cacaccaata cccgctgaga tccattacac actacttatg 10261 gatccaatga atgtttttta tcctggggaa tatgttggtt accttgtgga tgctgaaggt 10321 ggtgatatct atgggtcata ccagccaaca tacacatacg caattattgt gcaagaagtt 10381 gaaagagaag atgctgacaa tactagtttc ttaggaaaga tctatcagat cgatattggc 10441 tacaqtqaat ataaqataqt caqctctctt gatctgtaca agttctcaag gcctgatgaa 10501 ageteccaaa acagagacag tgeteccace acaccaacaa geeccaeega atteetgaet 10561 cctggtctga gaagcatccc tcctctttc tctggcaagg agagccacaa gtctccctcc 10621 accaaacacc attcccccag aaagctcaag gtgaatgctt taccagaaat cttaaaagaa 10681 gtgacatcag tggtggagca agcttggaag cttccagaat cagagcggaa aaagatcatt 10741 agacgettgt atttgaagtg geaccetgae aaaaateeag aaaateatga tattgetaat 10801 gaagtgttca agcacctgca gaatgaaatc aacagattag aaaaacaggc ttttctggat 10861 caaaatgcag acagagcttc aagaagaaca ttttcaacct ctgcatctcg atttcagtca 10921 gacaagtact catttcaaag attttacact tcgtggaatc aagaagccac aagtcataaa 10981 totgaaaggc aacagcaaag caaagagaaa tgccctcctt ctgctggaca gacatactct 11041 caaaggttct ttgttcctcc caccttcaag tcagtgggca atccagtgga agcccggaga 11101 tggttaagac aagccagagc aaacttctca gctgccagga atgaccttca caaaaatgcc 11161 aatgaatggg tgtgcttcaa gtgttacctt tccaccaagc tggctttgat tgcagccgac 11221 tatgctgtca gggggaaatc tgataaagat gtaaagccaa ctgcacttgc acaaaagata 11281 gaggagtaca gtcagcagct ggaaggactg acaaacgatg tgcacacatt ggaagcttat 11341 ggtgtagaca gcttgaaaac aaggtaccct gatttgcttc cttttccgca gattcccaat 11401 gacaggttca catctgaggt tgccatgagg gtgatggaat gcactgcctg tatcatcata 11461 aaacttgaaa attttataca acagaaggtg tga

Figure 8G



App No.: Inventors: 09/693,205

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LOCUS AF193556 12793 bp DNA PRI 07-FEB-2000 DEFINITION Homo sapiens sacsin (SACS) gene, complete cds. ACCESSION AF193556 VERSION AF193556.1 GI:6907041KEYWORDS . SOURCE human.

ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (bases 1 to 12793) AUTHORS Engert, J.C., Berube, P., Mercier, J., Dore, C., Lepage, P., Ge, B., Bouchard, J.P., Mathieu, J., Melancon, S.B., Schalling, M., Lander, E.S., Morgan, K., Hudson, T.J. and Richter, A. TITLE ARSACS, a spastic ataxia common in northeastern Quebec, is caused by mutations in a new gene encoding an 11.5-kb ORF JOURNAL Nat. Genet. 24 (2), 120-125 (2000) MEDLINE 20120709

REFERENCE 2 (bases 1 to 12793) AUTHORS Engert, J.C., Berube, P., Dore, C., Lepage, P., Ge, B., Hudson, T.J. and Richter, A. TITLE Direct Submission JOURNAL Submitted (08-OCT-1999) Genome Centre, Montreal General Hospital, 1650 Cedar Ave., Montreal, QC H3G 1A4, CanadaFEATURES Location/Qualifiers source 1..12793 /organism="Homo sapiens" /db xref="taxon:9606" /chromosome="13" /map="between D13S232 and D13S292" mRNA 1..12793 /gene="SACS" /product="sacsin" gene 1..12793 /gene="SACS" CDS 77..11566 /gene="SACS" /note="molecular chaperone" /codon start=1 /product="sacsin" /protein_id="AAF31262.1" /db_xref="GI:6907042" /translation="

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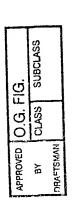
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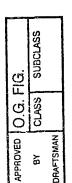
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